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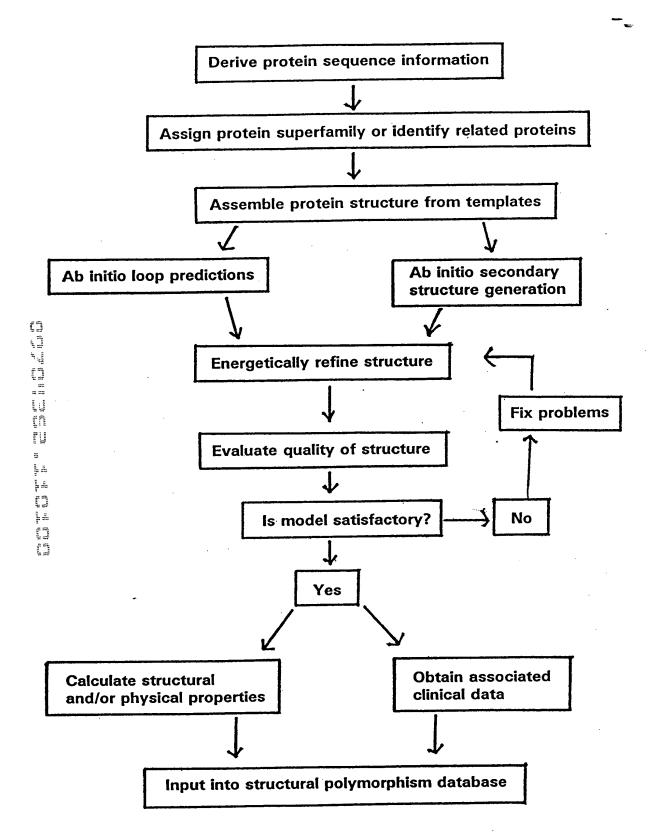
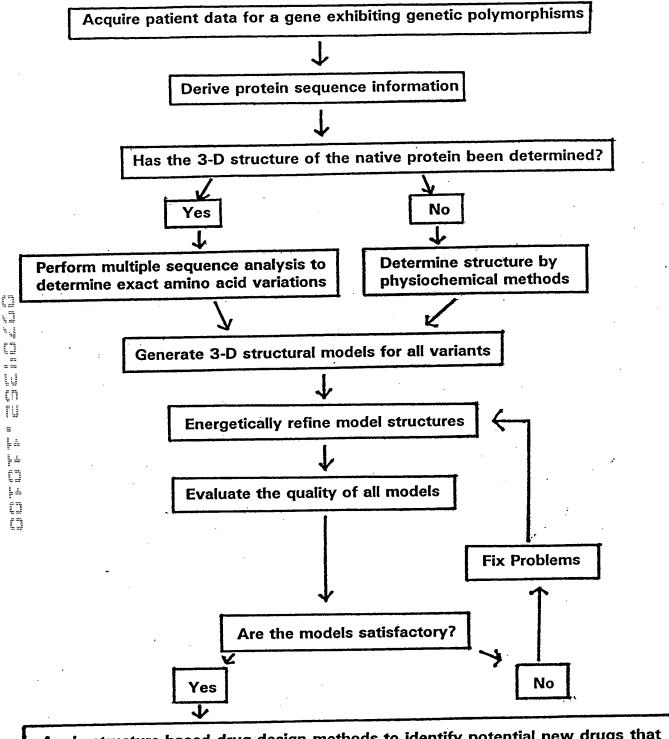


FIG. 1



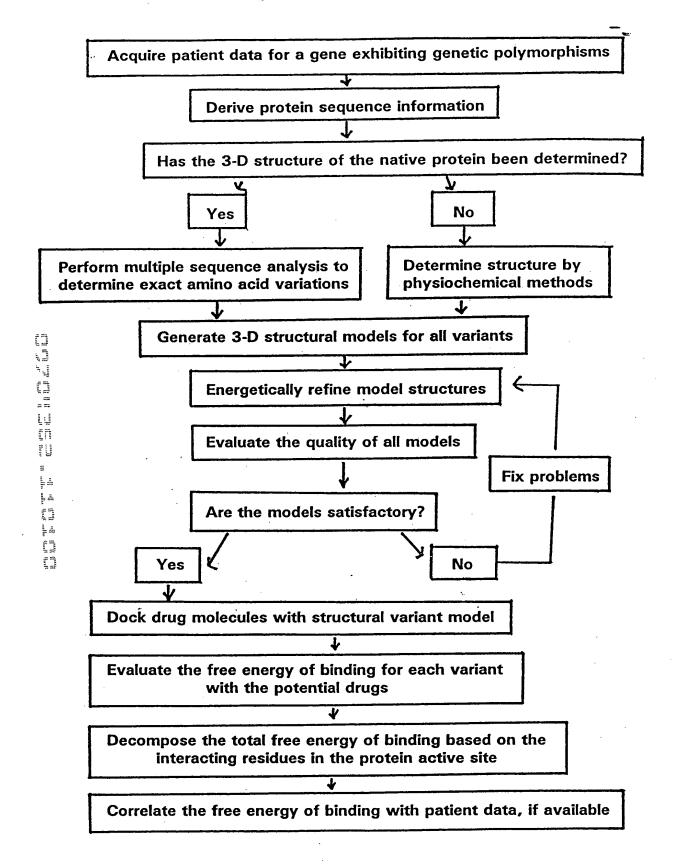


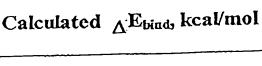


Apply structure-based drug design methods to identify potential new drugs that bind to the structural variant models

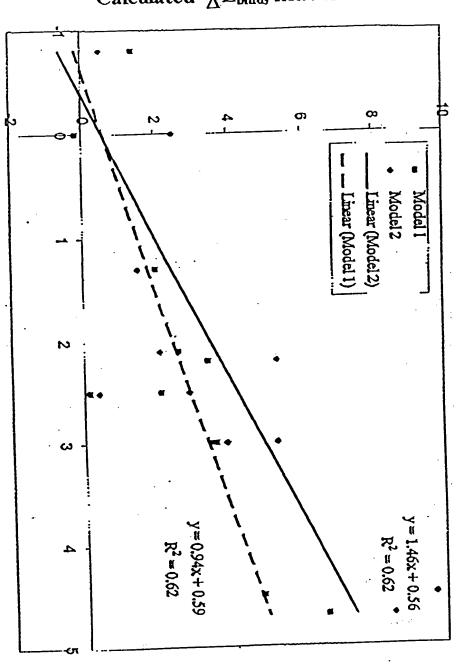






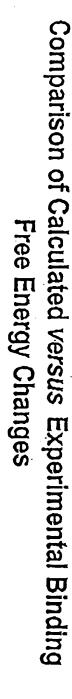


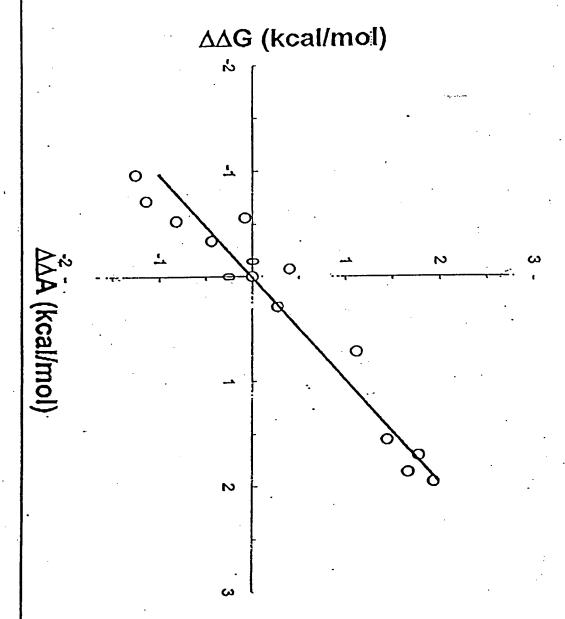
of Binding Energy upon Ligand Modifications in the Binding Correlation between Experimental and Calculated Changes Site of NS3



Expected AEbba kcal/mol

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FIG. J